AUG 2 5 2004 W

SEQUENCE LISTING

<110> Hammock, Bruce D.
 Kim, In-Hae
 Morisseau, Christophe
 Watanabe, Takaho
 Newman, John W.
 The Regents of the University of California

<120> Improved Inhibitors for the Soluble Epoxide Hydrolase

<130> 02307W-131010US

<140> US 10/817,334

<141> 2004-04-02

<150> US 60/460,559

<151> 2003-04-03

<160> 4

<170> PatentIn Ver. 2.1

<210> 1

<211> 555

<212> PRT

<213> Homo sapiens

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<223> human soluble epoxide hydrolase (sEH)

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Pro Leu Met Glu Glu Asn Cys Arg Lys Cys Ser Glu Thr Ala Lys Val 65 70 75 80

Cys Leu Pro Lys Asn Phe Ser Ile Lys Glu Ile Phe Asp Lys Ala Ile 85 90 95

Ser Ala Arg Lys Ile Asn Arg Pro Met Leu Gln Ala Ala Leu Met Leu 100 105 110

Arg Lys Cly Phe Thr Thr Ala Ile Leu Thr Asn Thr Trp Leu Asp

Asp Arg Ala Glu Arg Asp Gly Leu Ala Gln Leu Met Cys Glu Leu Lys 130 135 140

Met His Phe Asp Phe Leu Ile Glu Ser Cys Gln Val Gly Met Val Lys 145 150 155 160

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Phe Val Leu Val Pro Gln Met Ser Gln His Met Glu Asp Trp Ile Pro
500 505 510

His Leu Lys Arg Gly His Ile Glu Asp Cys Gly His Trp Thr Gln Met 515 520 525

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Asp Ala Arg Asn Pro Pro Val Val Ser Lys Met 545 550 555

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Pro Thr Glu Gln Leu Met Lys Gly Lys Ile Thr Phe Ser Gln Trp Val
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Ala Ala Arg Ser Ile Asn Arg Pro Met Leu Gln Ala Ala Ala Leu 100 105 110

Lys Lys Gly Phe Thr Thr Cys Ile Val Thr Asn Asn Trp Leu Asp 115 120 125

Asp Ser Asp Lys Arg Asp Ile Leu Ala Gln Met Met Cys Glu Leu Ser 130 135 140

Gln His Phe Asp Phe Leu Ile Glu Ser Cys Gln Val Gly Met Ile Lys 145 150 155 160

Pro Glu Pro Gln Ile Tyr Lys Phe Val Leu Asp Thr Leu Lys Ala Lys
165 170 175

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Leu Lys Arg Gly His Ile Glu Asp Cys Gly His Trp Thr Gln Ile Glu 515 520 525

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Ile Gln Asn Pro Ser Val Thr Ser Lys Ile 545 550

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Pro Thr Glu Gln Leu Met Lys Gly Lys Ile Thr Phe Ser Gln Trp Val 50 55 60

Pro Leu Met Asp Glu Ser Tyr Arg Lys Ser Ser Lys Ala Cys Gly Ala 65 70 75 80

Asn Leu Pro Glu Asn Phe Ser Ile Ser Gln Ile Phe Ser Gln Ala Met 85 90 95

Ala Ala Arg Ser Ile Asn Arg Pro Met Leu Gln Ala Ala Ile Ala Leu 100 105 110

Lys Lys Gly Phe Thr Thr Cys Ile Val Thr Asn Asn Trp Leu Asp 115 120 125

Asp Gly Asp Lys Arg Asp Ser Leu Ala Gln Met Met Cys Glu Leu Ser 130 140

Gln His Phe Asp Phe Leu Ile Glu Ser Cys Gln Val Gly Met Ile Lys 145 150 155 160

Pro Glu Pro Gln Ile Tyr Asn Phe Leu Leu Asp Thr Leu Lys Ala Lys
165 170 175

Pro Asn Glu Val Val Phe Leu Asp Asp Phe Gly Ser Asn Leu Lys Pro 180 185 190

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Ala Leu Arg Glu Leu Glu Lys Val Thr Gly Thr Gln Phe Pro Glu Ala 210 215 220

Pro Leu Pro Val Pro Cys Asn Pro Asn Asp Val Ser His Gly Tyr Val 230 235 Thr Val Lys Pro Gly Ile Arg Leu His Phe Val Glu Met Gly Ser Gly Pro Ala Leu Cys Leu Cys His Gly Phe Pro Glu Ser Trp Phe Ser Trp Arg Tyr Gln Ile Pro Ala Leu Ala Gln Ala Gly Phe Arg Val Leu Ala Ile Asp Met Lys Gly Tyr Gly Asp Ser Ser Ser Pro Pro Glu Ile Glu Glu Tyr Ala Met Glu Leu Leu Cys Lys Glu Met Val Thr Phe Leu Asp 315 310 Lys Leu Gly Ile Pro Gln Ala Val Phe Ile Gly His Asp Trp Ala Gly Val Met Val Trp Asn Met Ala Leu Phe Tyr Pro Glu Arg Val Arg Ala Val Ala Ser Leu Asn Thr Pro Phe Met Pro Pro Asp Pro Asp Val Ser Pro Met Lys Val Ile Arg Ser Ile Pro Val Phe Asn Tyr Gln Leu Tyr 375 Phe Gln Glu Pro Gly Val Ala Glu Ala Glu Leu Glu Lys Asn Met Ser 390 Arg Thr Phe Lys Ser Phe Phe Arg Ala Ser Asp Glu Thr Gly Phe Ile, 410 Ala Val His Lys Ala Thr Glu Ile Gly Gly Ile Leu Val Asn Thr Pro Glu Asp Pro Asn Leu Ser Lys Ile Thr Thr Glu Glu Glu Ile Glu Phe Tyr Ile Gln Gln Phe Lys Lys Thr Gly Phe Arg Gly Pro Leu Asn Trp 460 Tyr Arg Asn Thr Glu Arg Asn Trp Lys Trp Ser Cys Lys Gly Leu Gly Arg Lys Ile Leu Val Pro Ala Leu Met Val Thr Ala Glu Lys Asp Ile Val Leu Arg Pro Glu Met Ser Lys Asn Met Glu Lys Trp Ile Pro Phe Leu Lys Arg Gly His Ile Glu Asp Cys Gly His Trp Thr Gln Ile Glu

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<213> Mus musculus

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Met Asp Glu Ser Tyr Arg Lys Ser Ser Lys Ala Cys Gly Ala Asn Leu
50 55 60

Pro Glu Asn Phe Ser Ile Ser Gln Ile Phe Ser Gln Ala Met Ala Ala 65 70 75 80

Arg Ser Ile Asn Arg Pro Met Leu Gln Ala Ala Ile Ala Leu Lys Lys 85 90 95

Lys Gly Phe Thr Thr Cys Ile Val Thr Asn Asn Trp Leu Asp Asp Gly
100 105 110

Asp Lys Arg Asp Ser Leu Ala Gln Met Met Cys Glu Leu Ser Gln His 115 120 125

Phe Asp Phe Leu Ile Glu Ser Cys Gln Val Gly Met Ile Lys Pro Glu 130 135 140

Pro Gln Ile Tyr Asn Phe Leu Leu Asp Thr Leu Lys Ala Lys Pro Asn 145 150 155 160

Glu Val Val Phe Leu Asp Asp Phe Gly Ser Asn Leu Lys Pro Ala Arg 165 170 175

Asp Met Gly Met Val Thr Ile Leu Val His Asn Thr Ala Ser Ala Leu 180 185 190

Arg Glu Leu Glu Lys Val Thr Gly Thr Gln Phe Pro Glu Ala Pro Leu 195 200 205

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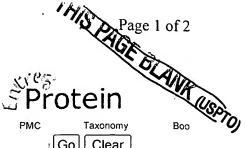
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PubMed Entrez Nucleotide Protein Genome Structure

PMC

Search Protein

for

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Clipboard Get Subsequence Details Feat

☐ 1: P80299. Soluble epoxide h...[gi:462371]

Limits

BLink, Domains, Links

P80299 LOCUS 554 aa linear ROD 15-JUN-2004

Soluble epoxide hydrolase (SEH) (Epoxide hydratase) (Cytosolic DEFINITION

epoxide hydrolase) (CEH).

ACCESSION P80299

P80299 GI:462371 VERSION

DBSOURCE swissprot: locus HYES RAT, accession P80299;

class: standard. created: Feb 1, 1994.

sequence updated: Feb 1, 1994. annotation updated: Jun 15, 2004.

xrefs: gi: 402631, gi: 402632, gi: 55929, gi: 55930, gi: 477003

xrefs (non-sequence databases): HSSPP34914, MEROPSS33.973, InterProIPR000073, InterProIPR003089, InterProIPR000639, InterProIPR006402, InterProIPR005833, InterProIPR005834, InterProIPR000379, PfamPF00561, PfamPF00702, PRINTSPR00111,

PRINTSPR00412, PRINTSPR00413, TIGRFAMsTIGR01509

Hydrolase; Peroxisome; Detoxification; Aromatic hydrocarbons KEYWORDS

catabolism; Direct protein sequencing.

SOURCE Rattus norvegicus (Norway rat)

ORGANISM Rattus norvegicus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

Rattus.

REFERENCE (residues 1 to 554)

AUTHORS Knehr, M., Thomas, H., Arand, M., Gebel, T., Zeller, H.D. and Oesch, F.

TITLE Isolation and characterization of a cDNA encoding rat liver cytosolic epoxide hydrolase and its functional expression in

Escherichia coli

JOURNAL J. Biol. Chem. 268 (23), 17623-17627 (1993)

MEDLINE 93352557 PUBMED 8349641

REMARK SEQUENCE FROM N.A.

STRAIN=Sprague-Dawley; TISSUE=Liver

REFERENCE 2 (residues 1 to 554)

AUTHORS Arand, M., Knehr, M., Thomas, H., Zeller, H.D. and Oesch, F.

TITLE An impaired peroxisomal targeting sequence leading to an unusual

bicompartmental distribution of cytosolic epoxide hydrolase

JOURNAL FEBS Lett. 294 (1-2), 19-22 (1991)

92077134 MEDLINE 1743286 **PUBMED**

REMARK SEQUENCE OF 450-554 FROM N.A., AND PARTIAL SEQUENCE.

TISSUE=Liver

COMMENT

This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. The original entry is available from http://www.expasy.ch/sprot

and http://www.ebi.ac.uk/sprot

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[FUNCTION] This enzyme acts on epoxides (alkene oxides, oxiranes)
            and arene oxides. Plays a role in xenobiotic metabolism by
            degrading potential toxic epoxides. Also determines steady-state
            levels of physiological mediators.
            [CATALYTIC ACTIVITY] An epoxide + H(2)O = a glycol.
            [SUBUNIT] Homodimer.
            [SUBCELLULAR LOCATION] Cytoplasmic and peroxisomal.
            [INDUCTION] By compounds that cause peroxisome proliferation such
            as clofibrate, tiadenol and fenofibrate.
            [SIMILARITY] Belongs to the AB hydrolase superfamily. Epoxide
            hydrolase family.
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11
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Aug 4 2004 12:36 34





Nucleotide

Protein Genome

Structure

History

STE BLANK BOLLSOTO, РМС Taxonomy Go Clear

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for

Clipboard Details Get Subsequence

☐ 1: JC4711. epoxide hydrolase...[gi:2135082]

BLink, Domains, Links

LOCUS JC4711 555 aa linear PRI 17-MAR-2000

epoxide hydrolase (EC 3.3.2.3) 2, cytosolic - human. DEFINITION

JC4711 ACCESSION

GI:2135082 VERSION JC4711 **DBSOURCE** pir: locus JC4711;

summary: #length 555 #molecular-weight 62615 #checksum 7933

genetic: #gene GDB:EPHX2 ##cross-references GDB:371845; OMIM:132811 #map position 8p21-8p12 #introns 34/2; 62/3; 116/1; 179/3; 220/3; 245/3; 277/3; 304/1; 315/3; 324/3; 353/2; 390/3; 414/3; 426/1;

460/3; 483/3; 510/3; 530/ 2

PIR dates: 16-Aug-1996 #sequence revision 16-Aug-1996 #text change

17-Mar-2000

KEYWORDS aromatic hydrocarbon catabolism; detoxification; ether hydrolase;

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

(residues 1 to 555) REFERENCE

AUTHORS Beetham, J.K., Tian, T. and Hammock, B.D.

TITLE cDNA cloning and expression of a soluble epoxide hydrolase from

human liver

JOURNAL Arch. Biochem. Biophys. 305 (1), 197-201 (1993)

MEDLINE 93343630 8342951 PUBMED

(residues 1 to 555) REFERENCE **AUTHORS** Sandberg, M. and Meijer, J.

Structural characterization of the human soluble epoxide hydrolase TITLE

gene (EPHX2)

JOURNAL Biochem. Biophys. Res. Commun. 221 (2), 333-339 (1996)

MEDLINE 96192049 **PUBMED** 8619856

This enzyme is involved in the conversion of harmful COMMENT

epoxide-containing compounds into diols.

FEATURES Location/Qualifiers

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> > /organism="Homo sapiens" /db xref="taxon:9606"

1..555 Protein

/product="epoxide hydrolase 2, cytosolic"

/EC number="3.3.2.3"

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121 iltntwlddr aerdglaqlm celkmhfdfl iescqvgmvk pepqiykfll dtlkaspsev

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